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65 70 75 80

Cys Ala Val Asn Thr Ile Lys Ser Asp Leu Glu Ile Phe Gly Arg Leu 85 90 95

Thr Pro Val Gly Asp Gly Ile Lys Met His Glu Thr Val Ile Asn Gln $100 \,$ $105 \,$ 110

Met Pro Pro Gln Ala Ala Val Ile Leu Leu Thr Asp Gly His Asn Asn 115 120 125

Leu Gly Met Asn Pro Val Glu Glu Val Lys Ser Ile Tyr Gln Thr Asn 130 135 140

Lys Ala Ile Ile Asp Gln Ile Val Ala Leu Asn Ser Gly Ser Val Leu 165 170 175

Val Asp Gly Leu Gln Leu Leu Gln Asn Pro Ala Val Cys Gln Glu Phe \$180\$

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gtt Val 65	gat Asp	ggt Gly	gct Ala	tct Ser	att Ile 70	gga Gly	aca Thr	gat Asp	ggt	aaa Lys 75	aat Asn	gtt Val	ata Ile	aag Lys	ctc Leu 80	240
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											cct Pro					480
											gta Val					528
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Leu Leu Gly Glu Ser Tyr Gly Asp Ile Glu Asp Val Arg Ala Gly Leu

624

195 200 205

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_	2 2			_	-		-			agt Ser		-			=	768
_	_		_			-	-	_		gga Gly	_		-		_	816
_			_				_	_		gat Asp			_	_		864
										ttt Phe						912
	-					-			-	aaa Lys 315	-	-		_		960
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Asn Asp Tyr Leu Gly Ala Thr Ala Phe Trp Met Arg Ala Leu His Lys 130 135 140

Asn Tyr Asp Ser Asn Tyr Gly Ile Ser Lys Leu Pro Asn Phe Lys Gly
145 150 155 160

Thr Thr Leu Asp Val Val Gly Leu Thr Ile Pro Val Thr Ile Ser Asp 165 170 175

Ile Lys Ile Ala Pro Trp Gly Met Phe Ala Phe Ala Gly Lys Lys Ser 180 185 190

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Leu Pro Ala Met Pro Ala Gly Phe Gly Tyr Ser Trp Gly Ala Gly Asn 210 215 220

Pro Phe Gly Asp Val Phe Pro Asn Lys Lys Arg Gly Asn Ala Trp Trp 225 230 235 240

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Val Asp Gly Ala Tyr Gly Arg Ala Asp Leu Gly Ser Leu Arg Asn Val 260 265 270

Val Ile Gly Asp Phe Leu Leu Asp Lys Ile Asp Leu Lys Arg Gln Gly 275 280 285

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Gly Val Ile Gly Trp Tyr Ala Ser Gly Asp Lys Val Asp Ser Arg Gly 305 310 315 320

Ala Ser Lys Arg Ile Pro Thr Leu Val Gly Asn Trp Ser Ala Thr Ser 325 330 335

Phe Gly Tyr Ser Gly Ala Tyr Gly Ile Gly Lys Asp Ser Val Phe Gly 340 345 350

Asn Thr Ile Ala Gly Ser Trp Gly Val Val Val Gln Leu Lys Asp Ile

4 (± y

355 360 365

Ser Phe Leu Glu Asn Leu Thr His Val Ile Arg Gly Ala Arg Ile Gln 370 375 380

Gly Thr Asn Asn Lys Asp Val Pro Glu His Leu Gly Leu Ser Tyr Val 385 390 395 400

Thr Thr Ile Tyr Asp Thr Arg Gly Gly Asp Asn Met Leu Tyr Leu Thr 405 410 415

Lys Lys Asp Tyr Ala Trp Glu Val Asp Phe Asp Thr Glu Tyr Lys Ile 420 425 430

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